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#6

1645

RAW SEQUENCE LISTING

DATE: 02/05/2002

PATENT APPLICATION: US/09/833,675B

TIME: 14:04:35

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02052002\I833675B.raw

ENTERED

2 <110> APPLICANT: Weimer, Thomas
W--> 3 <120> TITLE OF INVENTION: PROCESS FOR FINDING OLIGONUCLEOTIDE SEQUENCES
W--> 4 FOR NUCLEIC ACID AMPLIFICATION METHODS
6 <130> FILE REFERENCE: 06478.1454-00
W--> 7 <140> CURRENT APPLICATION NUMBER: 09/833,675B
8 <141> CURRENT FILING DATE: 2001-04-13
W--> 9 <160> NUMBER OF SEQ ID: 9
10 <170> SOFTWARE: PatentIn version 3.1
13 <210> SEQ ID NO: 1
14 <211> LENGTH: 27
15 <212> TYPE: DNA
16 <213> ORGANISM: Red Sea Bream Iridovirus
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19 <222> LOCATION: (1)..(27)
20 <223> OTHER INFORMATION: n=inosine
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26 <211> LENGTH: 18
27 <212> TYPE: DNA
28 <213> ORGANISM: Homo sapiens
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39 <223> OTHER INFORMATION: n=inosine
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45 <211> LENGTH: 21
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49 <221> NAME/KEY: misc_feature
50 <222> LOCATION: (1)..(21)
51 <223> OTHER INFORMATION: n=inosine
W--> 52 <400> SEQUENCE: 4

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53 agcacccctnt caggcagtag c 21
 56 <210> SEQ ID NO: 5
 57 <211> LENGTH: 20
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 59 <213> ORGANISM: Helianthus tuberosus
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 62 <222> LOCATION: (1)..(20)
 63 <223> OTHER INFORMATION: n=inosine
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 65 tgggtcncga aagnccttgt 20
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 71 <213> ORGANISM: Drosophila melanogaster
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 74 <222> LOCATION: (1)..(25)
 75 <223> OTHER INFORMATION: n=inosine
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 77 gctcatgntg cagcncctnc gagac 25
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 82 <212> TYPE: DNA
 83 <213> ORGANISM: Cyprinella galactura
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 89 catagntcac tccccgtgta 20
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RAW SEQUENCE LISTING

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Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02052002\I833675B.raw

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31

VERIFICATION SUMMARY

DATE: 02/05/2002

PATENT APPLICATION: US/09/833,675B

TIME: 14:04:36

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02052002\I833675B.raw

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L:7 M:283 W: Missing Blank Line separator, <140> field identifier
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L:17 M:283 W: Missing Blank Line separator, <220> field identifier
L:21 M:283 W: Missing Blank Line separator, <400> field identifier
L:22 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:29 M:283 W: Missing Blank Line separator, <400> field identifier
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L:41 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
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L:52 M:283 W: Missing Blank Line separator, <400> field identifier
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:60 M:283 W: Missing Blank Line separator, <220> field identifier
L:64 M:283 W: Missing Blank Line separator, <400> field identifier
L:65 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
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L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
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L:89 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
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L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:107 M:283 W: Missing Blank Line separator, <220> field identifier
L:111 M:283 W: Missing Blank Line separator, <400> field identifier
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9